RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information
Center (STIC) detected errors when processing the following computer readable
form:

Application Serial Number: _10/079,754_

Source: _OIPE_

Date Processed by STIC: _3/8/02_

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   APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A
   NOTICE TO COMPLY
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER
VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
        http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is
a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the
United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission
   User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
   Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
            Or
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,
   2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office,
   Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

O\PE

ERROR DETECTED   SUGGESTED CORRECTION   SERIAL NUMBER: 10|079,754

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 ____ Wrapped Nucleics
Wrapped Aminos

The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 ____ Invalid Line Length

The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 ____ Misaligned Amino
Numbering

The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4 ____ Non-ASCII

The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules.. Please ensure your subsequent submission is saved in ASCII text.

5 ____ Variable Length

Sequence(s) ____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 ____ PatentIn 2.0
"bug"

A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7 ____ Skipped Sequences
(OLD RULES)

Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8 ____ Skipped Sequences
(NEW RULES)

Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 ____ Use of n's or Xaa's
(NEW RULES)

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.

10 ____ Invalid <213>
Response

Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11 _√_ Use of <220>

Sequence(s) 13+14 missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12 ____ PatentIn 2.0
"bug"

Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 ____ Misuse of n

n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH – Biotechnology Systems Branch – 08/21/2001

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/079,754

DATE: 03/08/2002
TIME: 15:22:50

Errors on pp. 4&5

Input Set : A:\1068 SEQLIST.txt
Output Set: N:\CRF3\03082002\J079754.raw

```
   4 <110> APPLICANT: Glenn, Matthew
   5        Grigor, Murray R
   6        Molenaar, Adrian J
   7        Davis, Stephen R
   9 <120> TITLE OF INVENTION: Compositions isolated from bovine
  10        mammary gland and methods for their use.
  13 <130> FILE REFERENCE: 11000.1068
C--> 15 <140> CURRENT APPLICATION NUMBER: US/10/079,754
C--> 15 <141> CURRENT FILING DATE: 2002-02-19
  15 <150> PRIOR APPLICATION NUMBER: US 09/699,146
  16 <151> PRIOR FILING DATE: 2000-10-27
  18 <150> PRIOR APPLICATION NUMBER: US 60162,701
  19 <151> PRIOR FILING DATE: 1999-10-29
  21 <150> PRIOR APPLICATION NUMBER: US 09/644,190
  22 <151> PRIOR FILING DATE: 2000-08-22
  24 <150> PRIOR APPLICATION NUMBER: US 60/150,330
  25 <151> PRIOR FILING DATE: 1999-08-23
  27 <160> NUMBER OF SEQ ID NOS: 15
  29 <170> SOFTWARE: FastSEQ for Windows Version 4.0
  31 <210> SEQ ID NO: 1
  32 <211> LENGTH: 505
  33 <212> TYPE: DNA
  34 <213> ORGANISM: Bovine
  36 <400> SEQUENCE: 1
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  38 tgtcttcatt atggctctca tcctagccat gattagagct gattcatctg aagagaaacg   120
  39 tcacaggaaa cggaaaaaac atcatagagg atattttcaa caataccagc catatcaacg   180
  40 atatccacta aattatcctc ctgcgtatcc atttccttaa aatgctgctt agtaactaca   240
  41 ggacatgatt agagagattt ttcacaatga tttttcctac tctttctgtt gtgttgaaaa   300
  42 ccatctttca aatgaataaa acaaagaaaa aaaaatcagt caagtagttg cacaacacat   360
  43 acttggaatc aaatatcaat attttaaaac ataataatga tagtctctga actatgtaat   420
  44 tggtttctac tttcttttct ctgtcactta ccatgcatgc ttaataaatt gatctatcaa   480
  45 gcataaaaaa aaaaaaaaaa aaaaa                                          505
  47 <210> SEQ ID NO: 2
  48 <211> LENGTH: 585
  49 <212> TYPE: DNA
  50 <213> ORGANISM: Bovine
  52 <400> SEQUENCE: 2
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  54 actaaagaaa gaagcttgtc caagggaaaa ttaaaaagcc agcaattgaa gatgacatca   120
  55 ggtaccgctg tccaggactc caccaaatat gaagatcttt atctttatct tcattatggc   180
  56 tctcatccta gccatgatta gagctgattc atctgaagag aaacgtcaca ggaaacggaa   240
  57 aaaacatcat agaggatatt ttcaacaata ccagccatat caacgatatc cactaaatta   300
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Input Set : A:\1068 SEQLIST.txt
Output Set: N:\CRF3\03082002\J079754.raw

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58 tcctcctgcg tatccatttc cttaaaatgc tgcttagtaa ctacaggaca tgattagaga   360
59 gattttttcac aatgattttt cctactcttt ctgttgtgtt gaaaaccatc tttcaaatga   420
60 ataaaacaaa gaaaaaaaaa tcagtcaagt agttgcacaa cacatacttg gaatcaaata   480
61 tcaatatttt aaaacataat aatgacagtc tctgaactat gtaattggtt tctactttct   540
62 tttctctgtc acttaccatg catgcttaat aaattgatct atcaa                    585
64 <210> SEQ ID NO: 3
65 <211> LENGTH: 869
66 <212> TYPE: DNA
67 <213> ORGANISM: Bovine
69 <400> SEQUENCE: 3
70 gtccaaggga aaattaaaaa gccagcaatt gaagatgaca tcaggtaccg ctgtccagga   60
71 ctccaccaaa tatgaagatc tttatcttta tcttcattat ggctctcatc ctagccatga   120
72 ttagagctga ttcatctgaa gagaaacgtc acaggaaacg gaaaaaacat catgtatgta   180
73 ttcctctgat aatgtggtat agtataagaa tatttttttac tcagaatatt tattctataa   240
74 gagaatacat atttatcttt gaaatatatc tatacaatga ttagcttatg tgtccattga   300
75 attatctttt tatgatacac taggtaaaga ccccaaagac ttgtgtgctg ttactgttta   360
76 catagaaacc tataatgcca ccctgataaa gccagttatt ttctaagaaa agttatttct   420
77 gtttggtaaa taattgcctt cctctgaaga atactgaaat tctaattata ctaagtacaa   480
78 tatacaccta aaggaccata gaggctgcta aagaacatat attatgtggt cagtttccat   540
79 tttctgattt cttttgggag gtaatgtatg ttaattgacc aagttaaaga agagagacaa   600
80 tttcatttat ttaacatttt tcttaatgat atgtatcact ggtgtctcat attattttga   660
81 catgtctaat caggttgata ggtctccaga attcttacta atacagtaag tactgctact   720
82 gctactgcta agttgcttca gtcatgtcca actctgtgcg accccataga cagcagcgca   780
83 ccaggctccc ctgcccctgg gattctccag gcaaggatac tggagtgggt tgccatttca   840
84 gataaattca aatagcttca aaaaaaaaa                                      869
86 <210> SEQ ID NO: 4
87 <211> LENGTH: 604
88 <212> TYPE: DNA
89 <213> ORGANISM: Bovine
91 <400> SEQUENCE: 4
92 gaagtatttt cagttctata ataagatctc ataactgatg taattacaaa aacaaatgaa   60
93 ggatttcaag gtatttaaac acagcagttt tctagcaaag aacatctcct gaagcatcag   120
94 aatttcatct ttcatgactg gactccacca aatatgaaga tctttatctt tgtcttcatt   180
95 atggctctca tcctagccat gattagagct gattcatctg aagagaaacg tcacaggaaa   240
96 cggaaaaaac atcatagagg atattttcaa caataccagc catatcaacg atatccacta   300
97 aattatcctc ctgcgtatcc atttccttaa aatgctgctt agtaactaca ggacatgatt   360
98 agagagattt ttcacaatga ttttttcctac tctttctgtt gtgttgaaaa ccatctttca   420
99 aatgaataaa acaaagaaaa aaaaatcagt caagtagttg cacaacacat acttggaatc   480
100 aaatatcaat attttaaaac ataataatga tagtctctga actatgtaat tggtttctac   540
101 tttctttttct ctgtcactta ccatgcatgc ttaataaatt gatctatcaa gcataaaaaa   600
102 aaaa                                                                 604
104 <210> SEQ ID NO: 5
105 <211> LENGTH: 96
106 <212> TYPE: DNA
107 <213> ORGANISM: Bovine
109 <400> SEQUENCE: 5
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111 tcttaaaatg ctgcttacca actacaggac atgata                             96
113 <210> SEQ ID NO: 6
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114 <211> LENGTH: 525
115 <212> TYPE: DNA
116 <213> ORGANISM: Bovine
118 <400> SEQUENCE: 6
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120 gtcttcatta tggctctcat cctagccatg attagagctg attcatctga agagaaacgt       120
121 cacaggaaac ggaaaaaaca tcatgttgat aggtctccag aattcttact aatacaagag       180
122 gatattttca acaataccag ccatatcaac gatatccact aaattatcct cctgcgtatc       240
123 catttcctta aaatgctgct tagtaactac aggacatgat tagagagatt tttcacaatg       300
124 atttttccta ctctttctgt tgtgttgaaa accatctttc aaatgaataa aacaaagaaa       360
125 aaaaaatcag tcaagtagtt gcacaacaca tacttggaat caaatatcaa tattttaaaa       420
126 cataataatg atagtctctg aactatgtaa ttggtttcta ctttcttttc tctgtcactt       480
127 accatgcatg cttaataaat tgatctatca agcataaaaa aaaaa                       525
129 <210> SEQ ID NO: 7
130 <211> LENGTH: 58
131 <212> TYPE: PRT
132 <213> ORGANISM: Bovine
134 <400> SEQUENCE: 7
135 Met Lys Ile Phe Ile Phe Val Phe Ile Met Ala Leu Ile Leu Ala Met
136 1               5                   10                  15
137 Ile Arg Ala Asp Ser Ser Glu Glu Lys Arg His Arg Lys Arg Lys Lys
138             20                  25                  30
139 His His Arg Gly Tyr Phe Gln Gln Tyr Gln Pro Tyr Gln Arg Tyr Pro
140             35                  40                  45
141 Leu Asn Tyr Pro Pro Ala Tyr Pro Phe Pro
142     50                  55
144 <210> SEQ ID NO: 8
145 <211> LENGTH: 58
146 <212> TYPE: PRT
147 <213> ORGANISM: Bovine
149 <400> SEQUENCE: 8
150 Met Lys Ile Phe Ile Phe Ile Phe Ile Met Ala Leu Ile Leu Ala Met
151 1               5                   10                  15
152 Ile Arg Ala Asp Ser Ser Glu Glu Lys Arg His Arg Lys Arg Lys Lys
153             20                  25                  30
154 His His Arg Gly Tyr Phe Gln Gln Tyr Gln Pro Tyr Gln Arg Tyr Pro
155             35                  40                  45
156 Leu Asn Tyr Pro Pro Ala Tyr Pro Phe Pro
157     50                  55
159 <210> SEQ ID NO: 9
160 <211> LENGTH: 70
161 <212> TYPE: PRT
162 <213> ORGANISM: Bovine
164 <400> SEQUENCE: 9
165 Met Lys Ile Phe Ile Phe Ile Phe Ile Met Ala Leu Ile Leu Ala Met
166 1               5                   10                  15
167 Ile Arg Ala Asp Ser Ser Glu Glu Lys Arg His Arg Lys Arg Lys Lys
168             20                  25                  30
169 His His Val Cys Ile Pro Leu Ile Met Trp Tyr Ser Ile Arg Ile Phe
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RAW SEQUENCE LISTING   DATE: 03/08/2002
PATENT APPLICATION: US/10/079,754 TIME: 15:22:50

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             Input Set : A:\1068 SEQLIST.txt
             Output Set: N:\CRF3\03082002\J079754.raw

170          35                  40                  45
171 Phe Thr Gln Asn Ile Tyr Ser Ile Arg Glu Tyr Ile Phe Ile Phe Glu
172     50                  55                  60
173 Ile Tyr Leu Tyr Asn Asp
174 65                  70
176 <210> SEQ ID NO: 10
177 <211> LENGTH: 58
178 <212> TYPE: PRT
179 <213> ORGANISM: Bovine
181 <400> SEQUENCE: 10
182 Met Lys Ile Phe Ile Phe Val Phe Ile Met Ala Leu Ile Leu Ala Met
183 1               5                  10                  15
184 Ile Arg Ala Asp Ser Ser Glu Glu Lys Arg His Arg Lys Arg Lys Lys
185         20                  25                  30
186 His His Arg Gly Tyr Phe Gln Gln Tyr Gln Pro Tyr Gln Arg Tyr Pro
187         35                  40                  45
188 Leu Asn Tyr Pro Pro Ala Tyr Pro Phe Pro
189     50                  55
191 <210> SEQ ID NO: 11
192 <211> LENGTH: 21
193 <212> TYPE: PRT
194 <213> ORGANISM: Bovine
196 <400> SEQUENCE: 11
197 Phe Gln Gln Tyr Gln Pro Tyr Glu Arg Tyr Pro Leu Asn Tyr Pro Pro
198 1               5                  10                  15
199 Ala Tyr Pro Leu Ser
200             20
202 <210> SEQ ID NO: 12
203 <211> LENGTH: 59
204 <212> TYPE: PRT
205 <213> ORGANISM: Bovine
207 <400> SEQUENCE: 12
208 Met Lys Ile Phe Ile Phe Val Phe Ile Met Ala Leu Ile Leu Ala Met
209 1               5                  10                  15
210 Ile Arg Ala Asp Ser Ser Glu Glu Lys Arg His Arg Lys Arg Lys Lys
211         20                  25                  30
212 His His Val Asp Arg Ser Pro Glu Phe Leu Leu Ile Gln Glu Asp Ile
213         35                  40                  45
214 Phe Asn Asn Thr Ser His Ile Asn Asp Ile His
215     50                  55
217 <210> SEQ ID NO: 13
218 <211> LENGTH: 8
219 <212> TYPE: PRT
220 <213> ORGANISM: Artificail sequence
222 <220> FEATURE:
223 <223> OTHER INFORMATION: Artificial sequence
225 <221> NAME/KEY: VARIANT
226 <222> LOCATION: (1)...(8)
227 <223> OTHER INFORMATION: Xaa = Any Amino Acid
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C--> (handwritten annotation at line 220)

(handwritten) − unacceptable, must explain genetic source; see error summary sheet, item 11

```
    229 <400> SEQUENCE: 13
W--> 230 Xaa Xaa His His Xaa Xaa Tyr Xaa
    231  1                5
    233 <210> SEQ ID NO: 14
    234 <211> LENGTH: 8
    235 <212> TYPE: PRT
    236 <213> ORGANISM: Artificial Sequence
    238 <220> FEATURE:
    239 <223> OTHER INFORMATION: Artificial sequence        -Some error
    241 <400> SEQUENCE: 14
    242 Lys Lys His His Arg Lys Tyr Phe
    243  1                5
    245 <210> SEQ ID NO: 15
    246 <211> LENGTH: 267
    247 <212> TYPE: DNA
    248 <213> ORGANISM: Bovine
    250 <400> SEQUENCE: 15
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    252 tgtcttcatt atggctctca tcctagccat gattagagct gattcatctg aagagaaacg       120
    253 tcacaggaaa cggaaaaaac atcatagagg atattttcaa caataccagc catatcaacg       180
    254 atatccacta aattatcctc ctgcgtatcc atttccttaa aatgctgctt agtaactaca       240
    255 ggacatgatt agagagattt ttcacaa                                           267
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VERIFICATION SUMMARY
PATENT APPLICATION: US/10/079,754

DATE: 03/08/2002
TIME: 15:22:51

Input Set : A:\1068 SEQLIST.txt
Output Set: N:\CRF3\03082002\J079754.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:220 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13